TECH CENTER 1600/2900

· mandagh.

## SEQUENCE LISTING

1

- -110 Holms, Rupert D.
- +120 Regulatory/Unfolding Peptides of Ezrin
- +130. GJE-67
- +140+ 09/256,070
- -141- 2001-05-17
- +150+ FCT/GB00/03566
- +151. + 10000 09 15
- +1505 9921881.0 +151 999-09-17
- · 160: 28
- +170. FatentIn version 3.1
- + 310 + 1
- + 211 + 32
- -1112 FET
- +2130 Artificial Sequence
- 3.701
- + 223 + Hepreceptor peptide
- +400 + 1

Ala Arg Glu Glu Lys His Gln Lys Gln Leu Glu Arg Gln Gln Leu Glu

Thr Glu Lys Lys Arg Arg Glu Thr Val Glu Arg Glu Lys Glu Gln Met

- -0.210 + 0.00
- + .111.- 34
- HARLEY PET
- #213 Artificial Sequence
- 41220 ×
- +:223. Hepreceptor peptide
- -:20.
- <!d210 MISC\_FEATURE</pre>
- +12227 (14)..(14)
- (1223 Maa = Tyr(P) 아 7///
- -:400:- 3

Met Arg Glu Lys Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu

T:\Sequences\GJE\GJE-67.seq.ST25.txt/DNB/ehm

```
Lys Thr Lys Lys Ala Glu Arg Glu Leu Ser Glu Gln Ile Gln Arg Ala
Leu Gln
-1.110 - 3
·1.111 · 5
·LIP FPT
3213 · Artificial Sequence
(1)
-223 Hepredeptor peptide
44900 3
Thr Glu Lys Lys Arg
-1016 - 4
·::::: 9
\text{-1.111} + \text{--} \text{FFT}
Artificial Sequence
· Lin ·
Hepreceptor peptide
-:400.5 4
Thr Glu Lys Lys Arg Arg Glu Thr Val
10.10 · 5
10.11 · 11
Wille PPT
HILL: Artificial Sequence
(1
Hapreceptor peptide
+(4.00) + -5.
Thr Glu Lys Lys Arg Arg Glu Thr Val Glu Arg
               5
H210 - 6
·1211 · 5
```

<213 · Artificial Sequence

-1.111 · PF.T

3

```
-1320b
 Hepreceptor peptide
 -:4 10: -- 6
 Lys Lys Arg Arg Glu
 · 2100 7
 -111 8
 PRT
 +11: Artificial Sequence
Hepreceptor peptide
 -:4000 7
 Lys Lys Arg Arg Glu Thr Val Glu
 ·:1100 - 8
 · 111 · 10
 HIDIL FF.T
 Hols: Artificial Sequence
 \sim 10
 ·III Hepredeptor peptide
 <1400° - 8
 Lys Lys Arg Arg Glu Thr Val Glu Arg Glu
 -1715a 9
 <.11 11
 <212 > PRT
<213 > Artificial Sequence
 40000 s
 WIR Hepreceptor peptide
 +(4 (ii) + 9
 Lys Lys Arg Arg Glu Thr Val Glu Arg Glu Lys
 10.1105 10
 1211 - 12
 Hall PRT
 Mallie Artificial Sequence
```

4000 m

4

```
+22% Hepreceptor peptide
+14000 - 10
Lys Lys Arg Arg Glu Thr Val Glu Arg Glu Lys Glu
11.1101-11
-1111 8
PF.1
H2188 Artificial Sequence
Hepreceptor peptide
-14000-11
Lys Arg Arg Glu Thr Val Glu Arg
1.100 12
·1.11: 10
H. III PRT
H213 - Artificial Sequence
-1211G1
Hammadeptor peptide
-14000 12
Lys Arg Arg Glu Thr Val Glu Arg Glu Lys
<:113    Artificial Sequence</pre>
+:3255, +
Hammada Hepreseptor peptide
-(400 - 13)
Lys Arg Arg Glu Thr Val Glu Arg Glu Lys Glu
\pm 210 \cdot -14
-1.11.1 - 5
Hall: PET
3223 - Artificial Sequence
-1.2.2Ú -
223 · Hepreceptor peptide
```

T:\Sequences\GJE\GJE-67.seq.ST25.txt/DNB/ehm

```
-(400) 14
Arg Arg Glu Thr Val
+:210:- 15
+121.00 PF.T
3.133 Artificial Sequence
. . .
+...33 Hepreceptor peptide
-140ath 15
Arg Glu Thr Val Glu Arg Glu Lys Glu
\pm 1.1101 \pm 1.6
·1.:111: 5
·1.111 PF.T
+213 Artificial Sequence
1.1.10
+3.33 Hepreceptor peptide
+4000-16
Glu Arg Glu Lys Glu
-1.1100 - 17
1.11 - 14
+312 PPT
+3130 Artificial Sequence
-13<u>1</u>0 -
Fil. 3 - Hepreceptor peptide
-400-17
Glu Arg Glu Lys Glu Gln Met Met Arg Glu Lys Glu Glu Leu
-1210.- 18
-211 - 5
HILL: PET
213 - Artificial Sequence
-17.J

<
```

```
<400 € 18
Lys Glu Glu Leu Met
+12100 1 ·
+12110 1 ·
+12110 PFT
<!!!!! Artificial Sequence</pre>
-122015
Hander Hepreceptor peptide
-(4000) - 19
Lys Clu Glu Leu Met Leu Arg Leu Gln Asp Tyr Glu Glu
\cdot (-100) = 200
Rule: Artificial Sequence
111
Hills: Hepreceptor peptide
1000
HISC FEATURE
+0.00.08 - (11)...(11)
+0.12308 - Maa = Tyr(P)
(4:00) = 20
Lys Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu
1 5
-12111- 1.1
+21.55 PFT
Attificial Sequence
×:220 ×
Hopreceptor peptide
-:4002 21
Glu Glu Leu Met Leu Arg Leu Gln Asp Tyr Glu Glu
     Ξ,
0.110. L.
·:211. 1.
<21. · PFT
```

```
13 Artificial Sequence
0.120
4.2% Hepreceptor peptide
11.1200
MISC_FEATURE
COLL (10)..(10)
COLL Xaa = Tyr(F)
-(400)s 22
Glu Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu
-0.10. 23
-0.11. 11
+0.112 + PRT
Antificial Sequence
+1.23 - Hepreceptor peptide
+400+ 23
Gir Leu Met Leu Arg Leu Gln Asp Tyr Glu Glu
1 5.
02.10 · 24
· 111 · 11
FLI. PET
+.13+ Artificial Sequence
< 1.000 -
Hepreceptor peptide
H221 - MISC_FEATURE H222 - (9)..(9)
-0.013 \cdot \text{Maa} = \text{Tyr}(P)
4406 - 24
Glu Leu Met Leu Arg Leu Gln Asp Xaa Glu Glu
2010 - 25
· 21.. · PET
4.113 Artificial Sequence
-1220 -
```

```
4323> Hepreceptor peptide
- (40.0) - 25
Mot Leu Arg Leu Gln
-(210)- 26
-12111- 5
+211 > PET
H2130 Artificial Sequence
1. 05
Hepreceptor peptide
-,460 - 26
Gln Asp Tyr Glu Glu
+ 200 + 27
+201 + 5
+212 + PET
<!!! Artificial Sequence</pre>
+0.770 \pm
+1.13 - Hepreseptor peptide
e(1220 s
-2.11 - MISC_FEATURE
·222 · (3)..(3)
+ \text{Main} \cdot \text{Maa} = \text{Tyr}(P)
- 400 - 27
Glm Asp Kaa Glu Glu
+1210 - 28
4211 + 14
HILLS PRT
4013 - Artificial Sequence
-12:20 -
<223 - Hepreceptor peptide</p>
<400 - 28
Thr Glu Lys Lys Arg Arg Glu Thr Val Glu Arg Glu Lys Glu
```